SEQUENCE LISTING

<110> ITO, Yasuaki
FUJII, Ryo
HINUMA, Shuji
FUKUSUMI, Shoji
MARUYAMA, Minoru

<120> Novel Screening Method

<130> 3136 US0P

<140> US 10/542408
<141> 2005-07-15

<150> PCT/JP2004/000248
<151> 2004-01-15

<150> JP 2003-010001
<151> 2003-01-17

<150> JP 2003-104540
<151> 2003-04-08

<150> JP 2003-194497
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<150> JP 2003-329080
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<212> PRT
<213> Homo sapiens

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Leu Glu Gln Ala Asn Arg Thr Arg Phe Pro Phe Phe Ser Asp Val Lys
20 25 30
Gly Asp His Arg Leu Val Leu Ala Ala Val Glu Thr Thr Val Leu Val
35 40 45
Leu Ile Phe Ala Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
50 55 60
Val Ala Arg Arg Arg Arg Arg Gly Ala Thr Ala Cys Leu Val Leu Asn
65 70 75 80
Leu Phe Cys Ala Asp Leu Leu Phe Ile Ser Ala Ile Pro Leu Val Leu
85 90 95
Ala Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Ala Cys His
100 105 110
Leu Leu Phe Tyr Val Met Thr Leu Ser Gly Ser Val Thr Ile Leu Thr
115 120 125
Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val His Leu Gln
130 135 140
Arg Gly Val Arg Gly Pro Gly Arg Arg Ala Arg Ala Val Leu Leu Ala

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145                 150                 155                 160
Leu Ile Trp Gly Tyr Ser Ala Val Ala Ala Leu Pro Leu Cys Val Phe
                165                 170                 175
Phe Arg Val Val Pro Gln Arg Leu Pro Gly Ala Asp Gln Glu Ile Ser
            180                 185                 190
Ile Cys Thr Leu Ile Trp Pro Thr Ile Pro Gly Glu Ile Ser Trp Asp
        195                 200                 205
Val Ser Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
    210                 215                 220
Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
225                 230                 235                 240
Leu Thr Val Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
                245                 250                 255
Gln Gln Asp Phe Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
            260                 265                 270
Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu
        275                 280                 285
Ile Gln Asn Phe Lys Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
    290                 295                 300
Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
305                 310                 315                 320
Tyr Asn Met Thr Leu Cys Arg Asn Glu Trp Lys Lys Ile Phe Cys Cys
                325                 330                 335
Phe Trp Phe Pro Glu Lys Gly Ala Ile Leu Thr Asp Thr Ser Val Lys
            340                 345                 350
Arg Asn Asp Leu Ser Ile Ile Ser Gly
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<210> 2
<211> 1083
<212> DNA
<213> Homo sapiens

<400> 2

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gcggtggaga caaccgtgct ggtgctcatc tttgcagtgt cgctgctggg caacgtgtgc  180
gccctggtgc tggtggcgcg ccgacgacgc cgcggcgcga ctgcctgcct ggtactcaac  240
ctcttctgcg cggacctgct cttcatcagc gctatccctc tggtgctggc cgtgcgctgg  300
actgaggcct ggctgctggg ccccgttgcc tgccacctgc tcttctacgt gatgaccctg  360
agcggcagcg tcaccatcct cacgctggcc gcggtcagcc tggagcgcat ggtgtgcatc  420
gtgcacctgc agcgcggcgt gcggggtcct gggcggcggg cgcgggcagt gctgctggcg  480
ctcatctggg gctattcggc ggtcgccgct ctgcctctct gcgtcttctt ccgagtcgtc  540
ccgcaacggc tccccggcgc cgaccaggaa atttcgattt gcacactgat ttggcccacc  600
attcctggag agatctcgtg ggatgtctct tttgttactt tgaacttctt ggtgccagga  660
ctggtcattg tgatcagtta ctccaaaatt ttacagatca caaaggcatc aaggaagagg  720
ctcacggtaa gcctggccta ctcggagagc caccagatcc gcgtgtccca gcaggacttc  780
cggctcttcc gcaccctctt cctcctcatg gtctccttct tcatcatgtg gagccccatc  840
atcatcacca tcctcctcat cctgatccag aacttcaagc aagacctggt catctggccg  900
tccctcttct tctgggtggt ggccttcaca tttgctaatt cagcctaaa ccccatcctc  960
tacaacatga cactgtgcag gaatgagtgg aagaaaattt tttgctgctt ctggttccca 1020
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<211> 361
<212> PRT

<213> Mus musculus

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            20   25   30
Gly Asp His Arg Leu Val Leu Ser Val Val Glu Thr Thr Val Leu Gly
        35   40   45
Leu Ile Phe Val Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
    50   55   60
Val Ala Arg Arg Arg Arg Arg Gly Ala Thr Ala Ser Leu Val Leu Asn
65   70   75   80
Leu Phe Cys Ala Asp Leu Leu Phe Thr Ser Ala Ile Pro Leu Val Leu
                85   90   95
Val Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Val Cys His
            100   105   110
Leu Leu Phe Tyr Val Met Thr Met Ser Gly Ser Val Thr Ile Leu Thr
        115   120   125
Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val Arg Leu Arg
    130   135   140
Arg Gly Leu Ser Gly Pro Gly Arg Arg Thr Gln Ala Ala Leu Leu Ala
145   150   155   160
Phe Ile Trp Gly Tyr Ser Ala Leu Ala Ala Leu Pro Leu Cys Ile Leu
                165   170   175
Phe Arg Val Val Pro Gln Arg Leu Pro Gly Gly Asp Gln Glu Ile Pro
            180   185   190
Ile Cys Thr Leu Asp Trp Pro Asn Arg Ile Gly Glu Ile Ser Trp Asp
        195   200   205
Val Phe Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
    210   215   220
Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
225   230   235   240
Leu Thr Leu Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
                245   250   255
Gln Gln Asp Tyr Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
            260   265   270
Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu
        275   280   285
Ile Gln Asn Phe Arg Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
    290   295   300
Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
305   310   315   320
Tyr Asn Met Ser Leu Phe Arg Asn Glu Trp Arg Lys Ile Phe Cys Cys
                325   330   335
Phe Phe Phe Pro Glu Lys Gly Ala Ile Phe Thr Asp Thr Ser Val Arg
            340   345   350
Arg Asn Asp Leu Ser Val Ile Ser Ser
        355   360

<210> 4
<211> 1083
<212> DNA
<213> Mus musculus

<400> 4
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gtcgtggaga ccaccgttct ggggctcatc tttgtcgtct cactgctggg caacgtgtgt  180
gctctagtgc tggtggcgcg ccgtcggcgc cgtggggcga cagccagcct ggtgctcaac  240
ctcttctgcg cggatttgct cttcaccagc gccatccctc tagtgctcgt cgtgcgctgg  300
actgaggcct ggctgttggg gcccgtcgtc tgccacctgc tcttctacgt gatgacaatg  360
agcggcagcg tcacgatcct cacactggcc gcggtcagcc tggagcgcat ggtgtgcatc  420
gtgcgcctcc ggcgcggctt gagcggcccg gggcggcgga ctcaggcggc actgctggct  480
ttcatatggg gttactcggc gctcgccgcg ctgcccctct gcatcttgtt ccgcgtggtc  540
ccgcagcgcc ttcccggcgg ggaccaggaa attccgattt gcacattgga ttggcccaac  600
cgcataggag aaatctcatg ggatgtgttt tttgtgactt tgaacttcct ggtgccggga  660
ctggtcattg tgatcagtta ctccaaaatt ttacagatca cgaaagcatc gcggaagagg  720
cttacgctga gcttggcata ctctgagagc caccagatcc gagtgtccca acaagactac  780
cgactcttcc gcacgctctt cctgctcatg gtttccttct tcatcatgtg gagtcccatc  840
atcatcacca tcctcctcat cttgatccaa aacttccggc aggacctggt catctggcca  900
tccctttttct tctgggtggt ggccttcacg tttgccaact ctgccctaaa ccccatactg  960
tacaacatgt cgctgttcag gaacgaatgg aggaagattt tttgctgctt ctttttttcca 1020
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<213> Artificial Sequence

<220>
<223> primer

<400> 5
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<210> 6
<211> 20
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<220>
<223> primer

<400> 6
cgctgtggat gtctatttgc 20

<210> 7
<211> 30
<212> DNA
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<220>
<223> primer

<400> 7
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<210> 8
<211> 361
<212> PRT
<213> Rattus norvegicus

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Pro Asp Gln Val Asn Arg Thr His Phe Pro Phe Phe Ser Asp Val Lys
20 25 30
Gly Asp His Arg Leu Val Leu Ser Val Leu Glu Thr Thr Val Leu Gly
35 40 45
Leu Ile Phe Val Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
50 55 60
Val Val Arg Arg Arg Arg Arg Gly Ala Thr Val Ser Leu Val Leu Asn
65 70 75 80
Leu Phe Cys Ala Asp Leu Leu Phe Thr Ser Ala Ile Pro Leu Val Leu
85 90 95
Val Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Val Cys His
100 105 110
Leu Leu Phe Tyr Val Met Thr Met Ser Gly Ser Val Thr Ile Leu Thr
115 120 125
Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val Arg Leu Arg
130 135 140
Arg Gly Leu Ser Gly Pro Gly Arg Arg Thr Gln Ala Ala Leu Leu Ala
145 150 155 160
Phe Ile Trp Gly Tyr Ser Ala Leu Ala Ala Leu Pro Leu Cys Ile Leu
165 170 175
Phe Arg Val Val Pro Gln Arg Leu Pro Gly Gly Asp Gln Glu Ile Pro
180 185 190
Ile Cys Thr Leu Asp Trp Pro Asn Arg Ile Gly Glu Ile Ser Trp Asp
195 200 205
Val Phe Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
210 215 220
Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
225 230 235 240
Leu Thr Leu Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
245 250 255
Gln Gln Asp Tyr Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
260 265 270
Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu
275 280 285
Ile Gln Asn Phe Arg Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
290 295 300
Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
305 310 315 320
Tyr Asn Met Ser Leu Phe Arg Ser Glu Trp Arg Lys Ile Phe Cys Cys
325 330 335
Phe Phe Phe Pro Glu Lys Gly Ala Ile Phe Thr Glu Thr Ser Ile Arg
340 345 350
Arg Asn Asp Leu Ser Val Ile Ser Thr
355 360

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<212> DNA
<213> Rattus norvegicus

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gtcctggaga ccaccgttct gggactcatc tttgtggtct cactgctggg caacgtgtgt 180

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acc 1083

<210> 10
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<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 10
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<210> 11
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 11
cgctcctgaa cagcgacat 19

<210> 12
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> probe

<400> 12
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<210> 13
<211> 33
<212> DNA
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<220>
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<400> 13
gtcgacatgt cccctgagtg tgcgcagacg acg 33

<210> 14
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<212> DNA
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<220>
<223> primer

<400> 14
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<210> 15
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<212> DNA
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<220>
<223> primer

<400> 15
tccgagtgtc ccaacaagac tac 23

<210> 16
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<212> DNA
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<220>
<223> primer

<400> 16
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<210> 17
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<220>
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<400> 17
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<220>
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<400> 18

gtggtggcct tcacgtttg 19

<210> 19
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 19
cgctcctgaa cagcgacat 19

<210> 20
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<220>
<223> probe

<400> 20
caactccgcc ctaaacccca ttctgt 26

<210> 21
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_DNA
<222> (20)..(21)
<223> mixed DNA/RNA n stands for deoxy thymidine

<400> 21
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<210> 22
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_DNA
<222> (1)..(2)
<223> mixed DNA/RNA n stands for deoxy thymidine

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